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DATABASE BROWSING

EBI Dbfetch

```

ID  HPCCORBEA  standard; genomic RNA; VRL; 447 BP.
XX
AC  L39317;
XX
SV  L39317.1
XX
DT  30-JUN-1995 (Rel. 44, Created)
DT  21-OCT-2001 (Rel. 69, Last updated, Version 4)
XX
DE  Hepatitis C virus type 3 clone NL96 precursor protein gene, partial cds.
XX
KW  .
XX
OS  Hepatitis C virus type 3
OC  Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
OC  Hepacivirus.
XX
RN  [1]
RP  1-447
RX  MEDLINE; 95052487.
RX  PUBMED; 7525693.
RA  van Doorn L.J., Kleter B., Stuyver L., Maertens G., Brouwer H., Schalm S.,
RA  Heijntink R., Quint W.;
RT  "Analysis of hepatitis C virus genotypes by a line probe assay and
RT  correlation with antibody profiles";
RL  J. Hepatol. 21(1):122-129(1994).
XX
RN  [2]
RP  1-447
RX  MEDLINE; 97201609.
RX  PUBMED; 9049395.
RA  van Doorn L.J., Kleter G.E., Stuyver L., Maertens G., Brouwer J.T.,
RA  Schalm S.W., Heijntink R.A., Quint W.G.V.;
RT  "Sequence analysis of hepatitis C virus genotypes 1 to 5 reveals multiple
RT  novel subtypes in the Benelux countries";
RL  J. Gen. Virol. 76(Pt 7):1871-1876(1995).
XX
FH  Key          Location/Qualifiers
FH
FT  source       1..447
FT              /db_xref="taxon:40363"
FT              /mol_type="genomic RNA"
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FT              /db_xref="UniProt/TrEMBL:Q81286"
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FT      MMVGAATLCSALYVGDLGALFLVGQGFSWRHRQHWTVQDCNCISI"
FT      mat peptide      97..447
FT      /note="putative"
FT      /product="el protein"
XX
SQ      Sequence 447 BP; 82 A; 130 C; 114 G; 118 T; 3 other;
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      gctttgttct catgcttgct tacacccaca gccgggctgg agtaccgtaa tgcctccgga      120
      ctctacatgg taactaacga ctgcagtaac ggtagtatcg tgtatgaggc cggggatatt      180
      atcctccact tacctggctg tgtcccctgc gtacgctctg gcaatacatc aagatgctgg      240
      atccctgtga gcccyaaccg cgccgtgaag tcgccctgcg ccgccaccgc ctctctccgc      300
      acgcacgtgg atatgatggg gggrgcggcc accctatgct cagctctcta cgtaggagac      360
      ctttgtggag cgctatttct tgytyggcag gggttctcat ggagacatcg ccagcattgg      420
      actgtccagg actgcaactg ttccatc      447
//

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